STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

Source

Date Processed by STIC:

JFW0 02/16/2006

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS. PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER</u> <u>VERSION 4.4.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
 U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06

Raw Sequence Listing Error Summary

	15/6 25%
ERROR DETECTED	SUGGESTED CORRECTION SERIAL NUMBER: 10/567, 808
ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE	
Wrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
2Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.
3Misaligned Amino Numbering	The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
4Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
5Variable Length	Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <20>-<223> section that some may be missing.
6Patentin 2.0 "bug"	A "bug" in Patentin version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) Normally, Patentin would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
7Skipped Sequences (OLD RULES)	Sequence(s)missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped
•	Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
8Skipped Sequences (NEW RULES)	Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence. <110> sequence id number <400> sequence id number 000
9Use of n's or Xea's (NEW RULES)	Use of n's and/or Xna's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220><223> is MANDATORY if n's or Xna's are present. In <220> to <223> section, please explain location of a or Xnn, and which residue a or Xnn represents.
10Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
Use of <220>	Sequence(s)missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown," Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
12PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of Patentin version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
13 Misusc of n/Xaa	"n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid

AMC - Biotechnology Systems Branch - 09/09/2003



T DWA

RAW SEQUENCE LISTING DATE: 02/16/2006 PATENT APPLICATION: US/10/567,808 TIME: 12:55:34

Input Set : A:\08321-0136PC.TXT
Output Set: N:\CRF4\02162006\J567808.raw

```
4 <110> APPLICANT: Thomas Jefferson University
            Albert J. Wong
      7 <120> TITLE OF INVENTION: METHOD FOR RAPID IDENTIFICATION OF
              ALTERNATIVE SPLICING
     Ω
     10 <130> FILE REFERENCE: 08321-0136PC
C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/567,808
                                                               Does Not Comply
C--> 12 <141> CURRENT FILING DATE: 2006-02-08
                                                               Corrected Diskette Needed
     12 <150> PRIOR APPLICATION NUMBER: 60/493,759
     13 <151> PRIOR FILING DATE: 2003-08-08
     15 <160 - MEMBER OF SEQ ID NOS: S .
     17 <170> SOFTWARE: FastSEQ for Windows Version 4.0
     19 <210> SEQ ID NO: 1
     20 <211> LENGTH: 22
     21 <212> TYPE: DNA
     22 <213 > ORGANISM: Artificial Sequence
     24 <220> FEATURE:
     25 <223> OTHER INFORMATION: (selectable tag
     27 <400> SEQUENCE: 1
     28 gtcatgcata gcaattgtcg ac
     30 <210> SEQ ID NO: 2
     31 <211> LENGTH: 17
     32 <212> TYPE: DNA
     33 <213> ORGANISM: Artificial Sequence
     35 <220> FEATURE:
     36 <223> OTHER INFORMATION: (selectable tag
     38 <400> SEQUENCE: 2
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     39 tececegggg ggaateg
     41 <210> SEQ ID NO: 3
     42 <211> LENGTH: 27
     43 <212> TYPE: DNA
     44 <213> ORGANISM: Artificial Sequence
     46 <220> FEATURE:
     47 <223 > OTHER INFORMATION: (selectable tag
     49 <400> SEQUENCE: 3
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     50 atgcatagca acctcacgtg tgaatcg
     52 <210> SEQ ID NO: 4
     53 <211> LENGTH: 21
     54 <212> TYPE: DNA
     55 <213> ORGANISM: Artificial Sequence
     57 <220> FEATURE:
     58 <223 > OTHER INFORMATION affinity medium
     60 <400> SEQUENCE: 4
                                                                            21
     61 acacgtgagg ttgctatgca t
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RAW SEQUENCE LISTING

DATE: 02/16/2006

PATENT APPLICATION: US/10/567,808

TIME: 12:55:34

Input Set : A:\08321-0136PC.TXT

Output Set: N:\CRF4\02162006\J567808.raw

63 <210> SEQ ID NO: 5

64 <211> LENGTH: 58

65 <212> TYPE: DNA

66 <213> ORGANISM: Artificial Sequence

68 <220> FEATURE:

69 <223> OTHER INFORMATION: polynucleotide linking moiety

71 <400> SEQUENCE: 5

72 acacgatecg cagatgtecg cagttattee ttttttggaa taactgegga catetgeg · Same Rener

74 <210> SEQ ID NO: 6

75 <211> LENGTH: 30

76 <212> TYPE: DNA

77 <213> ORGANISM: Artificial Sequence

79 <220> FEATURE:

80 <223> OTHER INFORMATION: (arbitrary sequence

82 <400> SEQUENCE: 6

83 aaattootat goatgotoaa tggactgtgt

30

VERIFICATION SUMMARY

DATE: 02/16/2006

PATENT APPLICATION: US/10/567,808

TIME: 12:55:35

Input Set : A:\08321-0136FC.TXT

Output Set: N:\CRF4\02162006\J567808.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application No

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date